Appl. No. 09/700,712

Amendment dated: April 11, 2005

Reply to OA of: September 23, 2004

This listing of claims will replace all prior versions and listings of claims in the

application.

Listing of Claims:

Claims 1-16(cancelled).

17(currently amended). A Δ thy A strain of Vibrio cholerae deprived of its thy A

gene in the chromosome and thus tacing lacking the functionality of the thy A gene

comprising at least onel one episomal autonomously replicating DNA elements element

having a functional thy A gene that enables the strain to grow in the absence of thymine

in the growth medium and the one or several episomal autonomously replicating DNA

elements further comprising a structural [[gen]] gene encoding a homologous or

heterologous protein.

18(previously presented). A \triangle thy A strain of Vibrio cholerae wherein the strain

has been deprived of its thy A gene by site-directed mutagenesis in the V. Cholerae

chromosome for the deletion and/or insertion of nucleotides at the locus of the thy A

gene.

19(currently amended). The Δ thy A strain of Vibrio cholerae according to claim

17, wherein the at least one episomal autonomously replicating DNA element is a

plasmid.

20(currently amended). The Δ thy A strain of Vibrio cholerae according to claim

17, wherein the at least on [[the]] episomal autonomously replicating DNA element s

have element has a foreign thy A gene.

- 2 -

Appl. No. 09/700,712

Amendment dated: April 11, 2005 Reply to OA of: September 23, 2004

21(previously presented) . The Δ thy A strain of Vibrio cholerae according to claim 20, wherein the foreign thy A gene is an E. coli gene.

22(previously presented). The Δ *thy* A strain of *Vibrio cholerae* according to claim 17, wherein the encoded heterologous protein is selected from heat labile enterotoxin B-subunit of *Escherichia coli* (LTB) and *Schistosoma japonicum* glutathione S-transferase 26 kD protein (GST 26 kD).

23(previously presented). The Δ thy A strain according to claim 17, wherein the thy A gene of the chromosome has the nucleotide sequence SEQ ID NO: 1, or said nucleotide sequence which has some natural or unnatural nucleotide extensions, truncations, deletions or additions that do not interfere with the natural function of the nucleotide sequence.